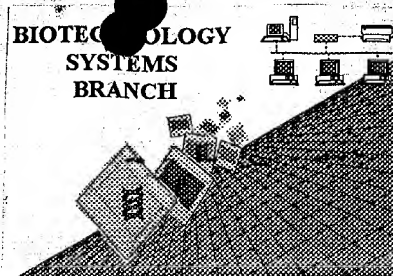


A. Nelson

## RAW SEQUENCE LISTING ERROR REPORT



File Copy  
#13

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/446,089  
Source: 1638  
Date Processed by STIC: 12/26/2000

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin30help@uspto.gov](mailto:patin30help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

### Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

09/446,089

insert hard return

~~SEQUENCE LISTING~~<110> SAKAKIBARA, Keiko ✓ FUKUI,  
Yuko ✓ TANAKA, Yoshikazu ✓ KUSUMI, Takaaki ✓ MIZUTANI, Masako ✓  
NAKAYAMA, Toru ✓<120> GENE ENCODING PROTEIN HAVING AURONE SYNTHESIZING  
ACTIVITY<130> 001560-377<140> US 09/446,089<141> 1999-12-17<150> ✓  
PCT/JP99/02045<151> 1999-04-16<150> JP 10/107296<151> 1998-04-17<160> 15 ✓  
<170> PatentIn version 3.0<210> 1<211> 1951<212> DNA<213> Antirrhinum  
majus<220><221> CDS<222> (96)..(1781)<400> 1  
↑ ↑ ↑ ↑

insert hard return at end of each response

The above is a sample of global format  
error. Please contact Robert Wax  
at 703-306-4119 for assistance.  
See cover sheet, too.

FYI

Please review the  
Sequence Listing to ensure that a corresponding explanation is presented in the <220> to  
<223> fields of each sequence which presents at least one n or Xaa.

1638

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/446,089  
 DATE: 12/26/2000  
 TIME: 13:36:45  
 Input Set : A:\PTO.txt  
 Output Set: N:\CRF3\12262000\I446089.raw

Does Not Comply  
 Corrected Diskette Needed

1 <110> APPLICANT: SAKAKIBARA, Keiko FUKUI,  
 2 Yuko TANAKA, Yoshikazu KUSUMI, Takaaki MIZUTANI, Masako  
 W--> 3 <120> TITLE OF INVENTION: GENE ENCODING PROTEIN HAVING AURONE SYNTHESIZING  
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 C--> 5 <141> CURRENT FILING DATE: 1999-12-17  
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 6 <170> SOFTWARE: PatentIn version 3.0<210> 1<211> 1951<212> DNA<213> Antirrhinum

ERRORED SEQUENCES

*See next page*

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VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/446,089

DATE: 12/26/2000  
TIME: 13:36:46

Input Set : A:\PTO.txt  
Output Set: N:\CRF3\12262000\I446089.raw

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L:4 M:283 W: Missing Blank Line separator, <130> field identifier  
L:5 M:270 C: Current Application Number differs, Replaced Current Application No  
L:5 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:0 M:282 W: Numeric Field Identifier Missing, <160> is required.  
L:7 M:283 W: Missing Blank Line separator, <220> field identifier  
L:7 M:256 W: Invalid Numeric Header Field, <220> has non-blank data  
L:328 M:256 W: Invalid Numeric Header Field, <220> has non-blank data  
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L:331 M:282 W: Numeric Field Identifier Missing, <212> is required.  
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L:408 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:0 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (0) Counted (6)

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